



**SEQUENCE LISTING**

<110> Evans, Ronald M.  
Forman, Barry M.

<120> SELECTIVE MODULATORS OF PEROXISOME  
PROLIFERATOR ACTIVATED RECEPTOR-GAMMA, AND METHODS FOR THE  
USE THEREOF

<130> SALK1470-2

<140> 09/155,252  
<141> 1998-09-21

<150> PCT/US96/05465  
<151> 1996-04-18

<160> 7

<170> FastSEQ for Windows Version 4.0

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tctgacagga gcctgtgaga ccaacagcct gacgggtctt cggtttaggg gacgcgggct 240
gagaagtac ac ttctgacag gactgtgtga cagacaagat ttgaaagaag cggtaacca 300
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att cca ttc aca aga gct gac cca atg gtt gct gat tac aaa tat gac 549  
 Ile Pro Phe Thr Arg Ala Asp Pro Met Val Ala Asp Tyr Lys Tyr Asp  
 55 60 65

ctg aag ctc caa gaa tac caa agt gcg atc aaa gta gaa cct gca tct 597  
 Leu Lys Leu Gln Glu Tyr Gln Ser Ala Ile Lys Val Glu Pro Ala Ser  
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cca cct tat tat tct gaa aag acc cag ctc tac aac agg cct cat gaa Pro Pro Tyr Tyr Ser Glu Lys Thr Gln Leu Tyr Asn Arg Pro His Glu 85 90 95	645
gaa cct tct aac tcc ctc atg gcc att gag tgc cga gtc tgt ggg gat Glu Pro Ser Asn Ser Leu Met Ala Ile Glu Cys Arg Val Cys Gly Asp 100 105 110	693
aaa gca tca ggc ttc cac tat gga gtt cat gct tgt gaa gga tgc aag Lys Ala Ser Gly Phe His Tyr Gly Val His Ala Cys Glu Gly Cys Lys 115 120 125 130	741
ggt ttt ttc cga aga acc atc cga ttg aag ctt att tat gat agg tgt Gly Phe Phe Arg Arg Thr Ile Arg Leu Lys Leu Ile Tyr Asp Arg Cys 135 140 145	789
gat ctt aac tgc cgg atc cac aaa aaa agt aga aat aaa tgt cag tac Asp Leu Asn Cys Arg Ile His Lys Lys Ser Arg Asn Lys Cys Gln Tyr 150 155 160	837
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agg ttt ggg cgg atg cca cag gcc gag aag gag aag ctg ttg gcg gag Arg Phe Gly Arg Met Pro Gln Ala Glu Lys Glu Lys Leu Leu Ala Glu 180 185 190	933
atc tcc agt gat atc gac cag ctg aac cca gag tct gct gat ctg cga Ile Ser Ser Asp Ile Asp Gln Leu Asn Pro Glu Ser Ala Asp Leu Arg 195 200 205 210	981
gcc ctg gca aag cat ttg tat gac tca tac ata aag tcc ttc ccg ctg Ala Leu Ala Lys His Leu Tyr Asp Ser Tyr Ile Lys Ser Phe Pro Leu 215 220 225	1029
acc aaa gcc aag gcg agg gcg atc ttg aca gga aag aca acg gac aaa Thr Lys Ala Lys Ala Arg Ala Ile Leu Thr Gly Lys Thr Thr Asp Lys 230 235 240	1077
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Tyr Asp Leu Lys Leu Gln Glu Tyr Gln Ser Ala Ile Lys Val Glu Pro  
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 His Glu Glu Pro Ser Asn Ser Leu Met Ala Ile Glu Cys Arg Val Cys  
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 Cys Lys Gly Phe Phe Arg Arg Thr Ile Arg Leu Lys Leu Ile Tyr Asp  
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 Arg Cys Asp Leu Asn Cys Arg Ile His Lys Lys Ser Arg Asn Lys Cys  
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 Ala Ile Arg Phe Gly Arg Met Pro Gln Ala Glu Lys Glu Lys Leu Leu  
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 Ala Glu Ile Ser Ser Asp Ile Asp Gln Leu Asn Pro Glu Ser Ala Asp  
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 Pro Leu Thr Lys Ala Lys Ala Arg Ala Ile Leu Thr Gly Lys Thr Thr  
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 Asp Lys Ser Pro Phe Val Ile Tyr Asp Met Asn Ser Leu Met Met Gly  
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 Glu Asp Lys Ile Lys Phe Lys His Ile Thr Pro Leu Gln Glu Gln Ser  
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50 55 60
Glu Gln Leu Phe Leu Leu Ile Phe Pro Arg Glu Asp Leu Asp Met Ile
65 70 75 80
Leu Lys Met Asp Ser Leu Gln Asp Ile Lys Ala Leu Leu Thr Gly Leu
85 90 95
Phe Val Gln Asp Asn Val Asn Lys Asp Ala Val Thr Asp Arg Leu Ala
100 105 110
Ser Val Glu Thr Asp Met Pro Leu Thr Leu Arg Gln His Arg Ile Ser
115 120 125
Ala Thr Ser Ser Ser Glu Glu Ser Ser Asn Lys Gly Gln Arg Gln Leu
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